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Richard, I., Roudaut, C., Fougerousse, F., Chiannilkuchai, N. and Beckmann, J. S. (1995). An STS map of the limb girdle muscular dystrophy type 2A region.

Mammalian Genome *6*, 754-756.

### **IN THE CLAIMS**

Kindly enter the following amended claims.

13

- 1. (2x Amended) An isolated nucleic acid sequence comprising:
  - a) the sequence represented in Figure 8 (SEQ ID NO:1-SEQ ID NO:4); or
  - b) the sequence represented in Figure 2 (SEQ ID NO:5, SEQ ID NO:68 and SEQ ID NO:69); or
  - c) a sequence obtained from a sequence defined in a) or b) by substitution, deletion or addition of one or more nucleotides with the proviso that said sequence still codes for said protease.
- 2. (Amended) An isolated nucleic acid sequence that is complementary to a nucleic acid sequence according to claim 1:
- 3. (Amended) A recombinant vector comprising in its structure a nucleotide sequence according to claim 1, under the control of regulatory elements, and involved in the expression of calpain activity in a LGMD2 disease.
- 4. (2x Amended) An isolated nucleic acid sequence encoding the amino acid sequence represented in Figure 2 (SEQ ID NO:6).
- 5. (Amended) An isolated amino acid sequence which is encoded by a nucleic acid sequence according to Claim 1, characterized in that it is a calcium dependent protease enzyme belonging to the calpain family, involved in the etiology of LGMD2.

### BECKMAN et al. - Appln. No. 08/836,734

6. (3x Amended) An isolated amino acid sequence according to claim 5 characterized in that either it contains the sequence such as represented in Figure 2 (SEQ ID NO:6), or the amino acid sequence of Figure 2 (SEQ ID NO:6) modified by deletion, insertion and/or replacement of one or more amino acids with the proviso that such amino acid sequence has the calpain activity involved in LGMD2 disease.



- 7. (Amended) An isolated amino acid sequence according to claim 5, characterized in that LGMD2 is LGMD2A.
- 8. (Amended) A host cell unable to express a calpain enzyme activity, characterized in that it is transformed or transfected with a nucleic acid sequence comprising the isolated nucleic acid sequence according to Claim 1.

414

- 12. (Amended) A method of screening, such method comprising the steps of:
  - providing an isolated amino acid sequence according to Claim 5 and
- determining ligands of said amino acid sequence, said ligands being selected from the group consisting of substrate(s), co-factors and regulatory components.
- 13. (Amended) A method of screening, such method comprising the steps of:
  - providing an isolated nucleic acid sequence according to Claim 1 and
- determining components which regulate expression of the novel calpain large subunit 1 (nCL1) gene.
- 14. (Amended) A method of screening, such method comprising the steps of:
  - providing an host cell according to claim 8 and
- determining components which regulate expression of the novel calpain large subunit 1 (nCL1) gene.
- 15. (2x Amended) A method for detecting an LGMD2 disease, such method comprising the steps of:

### BECKMAN et al. - Appln. No. 08/836,734

- selecting nucleotide sequences from one or more exons or flanking sequences of said one or more exons from an nCL1 gene,

- selecting primers specific for said one or more exons, or said flanking sequences, of said one or more exons,
- amplifying nucleic acid sequences of said one or more exons or said flanking sequences of one or more exons with said selected primers, and
- comparing the amplified sequence to the corresponding sequence obtained from Figure 2 (SEQ ID NO:5, SEQ ID NO:68 and SEQ ID NO:69) or Figure 8 (SEQ ID NO:1-SEQ ID NO:4) wherein a mutation in said amplified sequences is indicative of an LGMD2 disease.
- 16. (2x Amended) The method according to claim 15, characterized in that the primers are those selected from the group consisting of:
  - a) those described in Table 1 (SEQ ID NO:10-SEQ ID NO:17);
  - b) those described in Table 3 (SEQ ID NO:18-SEQ ID NO:67);
  - those including the introns-exons junctions of Table 2 (SEQ ID NO:71-SEQ ID NO:116); and
  - d) those derived from the primers in a), b), or c).

315

- 18. (2x Amended) A kit for the detection of a predisposition to LGMD2 by nucleic acid amplification characterized in that it comprises primers selected from the group consisting of:
  - a) those described in Table 1 (SEQ ID NO:10-SEQ ID NO:17);
  - b) those described in Table 3 (SEQ ID NO:18-SEQ ID NO:67);
  - those including the introns-exons junctions of Table 2 (SEQ ID NO:71-SEQ ID NO:116); and
  - d) those derived from the primers in a), b), or c).

20. (2x Amended) A pharmaceutical composition for the treatment of an LGMD2 disease characterized in that it contains a component selected from the group consisting of:

3/6

- a) an isolated nucleic acid sequence according to claim 1,
- b) a host cell according to claim 8, and
- c) an isolated amino acid sequence according to claim 5.

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22. (Amended) A method of screening a compound for its ability to modify the expression of the novel calpain large subunit 1 (nCL1) gene comprising contacting said compound with a host cell according to claim 8 and determining whether said compound modifies expression of said nCL1 gene in said host cell.

## **IN THE ABSTRACT**

Kindly enter the attached Abstract of the Disclosure.

# IN THE SEQUENCE LISTING

Kindly enter the attached paper and computer readable forms of the Sequence Listing in lieu of the Sequence Listing submitted on August 30, 2001.